



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/202,681

Source: 1652

Date Processed by STIC: 9/26/2000

RECEIVED

OCT 03 2000

TECH CENTER 1600/29

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
<http://www.uspto.gov/web/offices/pac/checker>

RECEIVED

2000

1600/2200

DATE: 09/26/2000
TIME: 16:15:49

```
Input Set : A:\sequence.txt
Output Set: N:\CRF3\092\2000\I202681.raw
```

Does Not Comply
Corrected Diskette Needed

SEQUENCE LISTING

1 (1) GENERAL INFORMATION:
2
3 (i) APPLICANT: recombinant Biocatalysis, Inc.
4
5 (ii) TITLE OF INVENTION: THERMOSTABLE PHOSPHATASES
6
7 (iii) NUMBER OF SEQUENCES: 54
8
9 (iv) CORRESPONDENCE ADDRESS:
10
11 (A) ADDRESSEE: Fish & Richardson P.C.
12
13 (B) STREET: 4025 Executive Square, Suite 1400
14
15 (C) CITY: La Jolla
16
17 (D) STATE: CA
18
19 (E) COUNTRY: USA
20
21 (F) ZIP: 92037
22
23 (v) COMPUTER READABLE FORM:
24
25 (A) MEDIUM TYPE: Diskette
26
27 (B) COMPUTER: IBM Compatible
28
29 (C) OPERATING SYSTEM: Windows 95
30
31 (D) SOFTWARE: FastSEQ for Windows Version 2.0b
32
33 (vi) CURRENT APPLICATION DATA:
34
35 (A) APPLICATION NUMBER: US 09/202,681
36
37 (B) FILING DATE: 18-Sep-2000
38
39 (C) CLASSIFICATION:
40
41 (vii) PRIOR APPLICATION DATA:
42
43 (A) APPLICATION NUMBER: 08/666,857
44
45 (B) FILING DATE: 19-JUN-1996
46
47 (A) APPLICATION NUMBER: 60/033,752
48
49 (B) FILING DATE: 19-JUN-1997
50
51 (viii) ATTORNEY/AGENT INFORMATION:
52
53 (A) NAME: Baile, Ph.D., LL.M.
54
55 (B) REGISTRATION NUMBER: 34,347
56
57 (C) REFERENCE/DOCKET NUMBER: 09010/015WO1
58
59 (ix) TELECOMMUNICATION INFORMATION:
60
61 (A) TELEPHONE: 619/678-5070
62
63 (B) TELEFAX: 619/678-5099
64
65 (C) TELEX:

ERRORS: SEQUENCES

210 (2) INFORMATION FOR SEQ ID NO: 14:
212 (i) SEQUENCE CHARACTERISTICS:
213 (A) LENGTH: 41 base pairs
214 (B) TYPE: nucleic acid
215 (C) STRANDEDNESS: single
216 (D) TOPOLOGY: linear
218 (ii) MOLECULE TYPE: cDNA
210 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
E--> 242 CCGAGGACTA AACATAGTCT AAGTAATTAG C

31

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/202,681

DATE: 09/26/2000

TIME: 16:15:51

Input Set : A:\sequence.txt

Output Set: N:\CRF3\09262000\I202681.raw

L:27 M:220 C: Keyword misspelled or invalid format. [(A) APPLICATION NUMBER:]
L:28 M:220 C: Keyword misspelled or invalid format. [(B) FILING DATE:]
L:242 M:204 E: No. of Bases differ, LENGTH:Input:34 Counted:31 SEQ:14